



Results 1 - 9 of 9 (0.10 sec)

$$[2\pi i \operatorname{Im}(\alpha)] \otimes [2\pi i \operatorname{Im}(\beta)] = A[1] \otimes B[1] \cong A \otimes B[2]$$

INTERNAL: from russpda.org

CA Klein, M Wilcz, J Pote, C Vermeulen. The Journal of ... 2002 - jupn.cupress.org
... HA-1, CD45, and PBGD expression were quantified in all test samples using these calibration
functions. ... The isolated DNA was subjected to whole genome amplification and CGH. ... DNA
methylation has been proposed as the primary mechanism for the inactivation of genes (24). ...
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... Date: Nov. 22 2007 (54) MULTIPLEXED METHYLATION DETECTION METHODS (76) Inventor: Jian-Bing Fan, San Diego, CA (US) Correspondence Address: MCDERMOTT, WILL & EMERY
4370 LA JOLLA VILLAGE DRIVE, SUITE 700 SAN DIEGO, CA 92122 (US) (21) Appl. ...

... The invention also provides a **calibration** curve comprising a first axis indicating the amount of unmethylated template population compared to methylated template ... The invention methods for detecting **methylation** allow for **whole genome amplification** after bisulfite ...

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147721,1 from clinchem.org

... To address these limitations, we have combined **whole genome amplification** by primer extension ... the "competitive" nature of the amplification of the target and quantitation **standard**. ... Regulatory region mutations as well as **methylation** represent additional possible mechanisms ...

58 Bioscience - US Patent App. 2010/0193,957, 2008 - freepatentsonline.com
... addition, preparation of target nucleic acids can be accomplished using **whole genome
amplification** techniques (eg. ... also be used to estimate the average amount of **methylation** for one ...
by including nucleotide analogs having higher binding affinities than **standard** nucleotides for ...

... of target nucleic acids can be accomplished using **whole genome amplification** techniques (eg, Lizardi, US Pat. No. 6,124,120). [0041] In some embodiments, target nucleic acids are chemically treated prior to analysis. For example, analysis of the **methylation** state of cytosines ...

H Schroeder, AC Humbert, V Kozel, D Gieser . - Experimental .., 1985 - Elsevier
Cited by 16 - Standard journal - Pl. Librai - All 3 versions

US Patent App. 2006/... 2006 - freepatentonline.com
... labeling reagents, which relied on stable isotopes, which are expensive, and not flexible to differential labeling of more than two mixtures of peptides; labeling methods limited only to **methylation** of carboxy ... **Standard** IUPAC nomenclature is used for describing pooled nucleotides ...
Patented sequence - Cysteine

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methylation ("whole genome amplification") Search

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